

Figure 1 (SEQ ID NO 1): Amino acid sequence of the protein PPTA from *P. chrysogenum* coded by the nucleic acid molecule according to the invention (illustrated from the N-terminus to the C-terminus).

1	MVDPSVSGIT	KMDTNDIKQN	DIPKDQPTLV	RWYMDVRRWD	EKYFDLPLLE
51	TLTQPDQAAV	KKYYQTSDKR	LSLASQLLKY	YYIHQATGTP	WSKIEIQ RTP
101	MPENRPFYDS	SLDFNVSHQA	GLTLFAGTRA	ATAHSLSGGP	QTLPRVGIDV
151	ACVDEPSRRR	ANRPPKTLAD	LATFVDVFSD	VLSLRELATI	KNPYATLKLA
201	RELGLNKSDP	SKDDQEV LAA	YGIRLFYSIW	ALKEAYLKMT	GDGLLASWIK
251	DLEFTNVVPP	EPVQTVGFAG	DPSATHAPSV	QNWGRPYS DV	KISLRGIPDH
301	SVRVQPVGFE	SDYIVATAAS	GPNIGSVSRQ	VVNDS DHHL	PGRITAFDSE
351	TGLQNVRI PP	IALRSIGDGD	PWRVDSKISD	PWLPMQEVDI	EIDIRPCADG
401	RCEHLRDLPS	F			

Figure 2 (SEQ ID NO 2): Genomic DNA sequence of the coding region of the pptA gene of *P. chrysogenum* from the translation start codon (ATG) to the translation stop codon (TAA). The intron is underscored. A single strand from the 5'- to 3'-direction is illustrated.

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1  atggtagacc ccagtgtgtc tgggaattgtg agtagccaca tagcctccat
51 gagtgcaccc actgaccaat ttcagaccaa aatggatacc aatgatatca
101 aacagaatga catccccaag gaccagccca cgttggtccg atggtacatg
151 gatgtcagac gttgggatga aaaatacttt gatctccctt tgcttgaaac
201 cttaacacag cctgatcagg cagctgtcaa gaagtactat caaacatcgg
251 acaagcgcct gtccttggcc tcccagttgc tgaaatatta ctacattcac
301 caagccactg gcactccctg gagcaagatt gagatccagc gtactccgat
351 gcccgaaaat cgaccattct acgattcaag cctggatttc aacgtcagcc
401 atcaggctgg tctcactctg ttgcgaggca cgcgtgccgc aacagcccac
451 tccttatccg gtggacctca aacattgcct cgcgtgggaa ttgacgttgc
501 gtgtgttgat gaaccctctc gtcgtcgtgc taatcgtccc ccgaagacac
551 ttgccgacct tgcaaccctc gtggatgtct tcagtgacgt tctctcactc
601 cgtgagcttg cgaccatcaa gaaccgcgtac gcgactctta aattggctcg
651 tgagcttggt ctgaataaaa gtgaccgcgag caaagacgac caggaagtcc
701 ttgctgccta cggcattcgg ctgttctact cgatttgggc tctcaaggag
751 gcttacttga aaatgaccgg agacggcctt ctggcctctt ggataaagga
801 tctggaattc acaaacgttg ttccccccga accagttcaa acagtcggat
851 ttgctggtga tccttctgcc actcacgcgc ctcgggtcca aaattggggc
901 cggccttact ccgatgtcaa aatctccttg cgtggcattc ctgaccattc
951 tgtgcgcggt cagctcgtcg gcttcgagtc cgactacata gttgccacgg
1001 ccgcgtcggg ccccaatatt ggatccggtt cgcggcaggt agtcgtgaat
1051 gacagcgatc accatctgcc agggcgatc acagccttcg actctgagac
1101 tggactccag aacgtccgca ttcccccaat cgcgcttcga tcaattggcg
1151 atggggaccc ctggcgtgtg gactcgaaaa tcagcgaccc ctggctcccc
1201 atgcaggagg tcgatattga aatcgatatc cggccctgtg cggatggtcg
1251 ttgcgagcac ctacgggatt taccaagcct ttaa

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Figure 3 (SEQ ID NO 3): cDNA sequence of the coding region of the pptA gene of *P.*

chrysogenum from the translation start codon (ATG) to the translation stop codon (TAA); a single strand from the 5'- to 3'-direction is illustrated.

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1   atggtagacc ccagtgtgtc tggaattacc aaaatggata ccaatgatat
51  caaacagaat gacatcccca aggaccagcc cacgttggtc cgatggtaca
101 tggatgtcag acgttgggat gaaaaatact ttgatctccc ttgcttgaa
151 accttaacac agcctgatca ggcagctgtc aagaagtact atcaaacatc
201 ggacaagcgc ctgtccttgg cctcccagtt gctgaaatat tactacattc
251 accaagccac tggcactccc tggagcaaga ttgagatcca gcgtactccg
301 atgcccgaaa atcgaccatt ctacgattca agcctggatt tcaacgtcag
351 ccatcaggct ggtctcactc tgttcgcagg cacgcgtgcc gcaacagccc
401 actccttata cgggtggacct caaacattgc ctgcgctggg aattgacggt
451 gcgtgtgttg atgaaccctc tcgtcgtcgt gctaatacgt ccccgagac
501 acttgccgac cttgcaacct tcgtggatgt cttcagtgac gttctctcac
551 tccgtgagct tgcgaccatc aagaaccctg acgcgactct taaattggct
601 cgtgagcttg gtctgaataa aagtgaccctg agcaaagacg accaggaagt
651 ccttgctgcc tacggcattc ggctgttcta ctcgatttgg gctctcaagg
701 aggcttactt gaaaatgacc ggagacggcc ttctggcctc ttggataaag
751 gatctggaat tcacaaacgt tgttcccccc gaaccagttc aaacagtcgg
801 atttgctggt gatccttctg ccactcacgc gccctcggtc caaaattggg
851 gccggcctta ctccgatgtc aaaatctcct tgcgtggcat tcctgaccat
901 tctgtgcgcg ttcagcccgt cggcttcgag tccgactaca tagttgccac
951 ggccgcgtcg ggccccaata ttggatccgt ttgcgcggcag gtagtcgtga
1001 atgacagcga tcaccatctg ccagggcgta tcacagcctt cgactctgag
1051 actggactcc agaacgtccg cattccccca atcgcgcttc gatcaattgg
1101 cgatggggac ccctggcggtg tggactcgaa aatcagcgac ccctggctcc
1151 ccatgcagga ggtcgatatt gaaatcgata tccggccctg tgcggatggt
1201 cgttgcgagc acctacggga tttaccaagc ttttaa

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Figure 4 (SEQ ID NO 4): Genomic DNA sequence of a Sall fragment of a genomic clone of the pptA gene (a single strand from the 5'- to 3'-direction is illustrated). The translation start codon (ATG) and the translation stop codon (TAA) of the coding region are underscored and illustrated in bold; the intron is underscored.

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1   gtcgaccgaa gtgggtttcgg ttcactcgca catcaagacc accgatcagc
51  tcttgcccg ccttctttgt cttgttggca gactcggcaa gcaaaatgag
101 cccggcgcat gtacccacg tccgtttgcg atccactctg cataaccac
151 gtattagatc gaattgatat ggactaaccg ggttcactca ctttacgaat
201 tctcgcagtg gctcgagaag atttgacctt gctgcgacta aagacatagt
251 ggtactctcg cctccgggca agaccaggcc gtcgcatgtt gccagttctt
301 gtggcgctcg tacttcaatg aagtgccatt ccgacggctg cgcttgctca
351 gcggcctttt tcaaaagctg cacatgctca aagaatgcgc cctgtagggc
401 caggactcca acagtgatag ccatttcctc tgaagatcgg aattgcggac
451 cctccgagct cgggtgcttc ttgatattga tgactctttt taaagcacat
501 gactttgact ttccggcggg gaacgtatca acacgtgatg gcggcctatc
551 tccatcttta attccacgcg acatcaggat atcgtgagag ctctcggacg
601 attcctgcgc actttgaaaa cagactgcat aaccgaggca ttatagtata
651 aaacaaatag actcacctac agaaagagtg ataagttagg tcctatacct
701 gtttccaatg tttctctctc ttgctggatc agctttaaca tatctatgga
751 tggatatctt gatagtcata gtcatattgc gcttgctatt gcatgtctct
801 ttgctacatc ctatttatgg tattatgtac acggcctgtt tctcgtttgc
851 cggcctattg atgtatacat gtattgggtg aggtagttaa tgccctcgct
901 tatcgacacg tgctgataga taaggacccc gataagacgc caacatggct
951 tctatccagg tgtggatgct ccgcatccaa ggtgcgaata tacgagatca
1001 caatgcaatg gttagaccca gtgtgtctgg aattgtgagt agccacatag
1051 cctccatgag tgcacccact gaccaatttc agaccaaagt ggataccaat
1101 gatatacaac agaattgacat cccaaggac cagcccacgt tggccgatg
1151 gtacatggat gtcagacgtt gggatgaaaa atactttgat ctccctttgc
1201 ttgaaacctt aacacagcct gatcaggcag ctgtcaagaa gtactatcaa
1251 acatcggaca agcgcctgtc cttggcctcc cagttgctga aatattacta
1301 cattcaccaa gccactggca ctccctggag caagattgag atccagcgta
1351 ctccgatgcc cgaaaatcga ccattctacg attcaagcct ggatttcaac
1401 gtcagccatc aggttggtct cactctgttc gcaggcacgc gtgcgcgaac
1451 agcccactcc ttatccggtg gacctcaaac attgcctcgc gtgggaattg
1501 acgttgcggtg tgttgatgaa ccctctcgtc gtcgtgctaa tcgtcccccg
1551 aagacacttg ccgaccttgc aaccttcgtg gatgtcttca gtgacgttct
1601 ctcactccgt gagcttgcca ccataagaa cccgtacgcg actcttaa
1651 tggctcgtga gcttggtctg aataaaaagt acccgagcaa agacgaccag
1701 gaagtccttg ctgcctacgg cattcggctg ttctactcga tttgggctct
1751 caaggaggct tacttgaaaa tgaccggaga cggccttctg gcctcttgga
1801 taaaggatct ggaattcaca aacgttggtc ccccgaacc agttcaaaca
1851 gtcggatttg ctggtgatcc ttctgccact cacgcgcctc cgggtccaaa
1901 ttggggccgg ccttactcgc atgtcaaaat ctecttgctg ggcattcctg
1951 accattctgt gcgcgttcag ctcgctcggc tcgagtcoga ctacatagtt
2001 gccacggccg cgtcggggcc caatattgga tccgtttcgc ggcaggtagt
2051 cgtgaatgac agcgatcacc atctgccagg gcgtatcaca gccttcgact
2101 ctgagactgg actccagaac gtccgcattc cccaatcgc gcttcgatca
2151 attggcgatg gggacccctg gcgtgtggac tcgaaaatca gcgacccctg
2201 gctcccatg caggaggtcg atattgaaat cgatatccgg ccctgtgcgg

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2251 atggtcgttg cgagcaccta cgggattttac caagcttttta aattccttct
2301 tgctgggata tgaccaggcg accatgcacc cgagttatTTT gcataattgca
2351 tctcctcatc tcatattcct ttctgagcgt gtttttcgga gcgataatta
2401 cccttgaaca tatttctgca ttgctgtatt gccattagcg aaaattcccg
2451 agctagttgt agttgatttc ctggaacgct gggggagtgC cgctcagatg
2501 ttcattctcca ataagccccT caatgaatct tcaattcatc ggatccaagg
2551 tcaatcttcg agatcaagtg caagttgccc agaaagcacg ggtaaagaaa
2601 ccaagcctat ttctattcta tggctctaatg taaactaaaa atgtagaagg
2651 aagaaaagca agtatccaac agtaggcggg tcatgacatg cgtgtgCgct
2701 aaggatatat acatttcgaa ttgcaaagag ggaagaggTg aatcaggagt
2751 gaaatgtgtg tcaagaggca atgtcaatgt caagatcatt gttgctctca
2801 tgagcagtca cggattgtgt cggattgttc ggcgtctggg gccctcagat
2851 tctatttctg ggtcatgagc ttgagagtag gtaccgaaga agtgagcagt
2901 attatactgc agtgagtgtt taggggggaat tccttctggT gaattgtggc
2951 gttcgggggt gctctccggT cttatgggtc ttaatctgga tgcccgatag
3001 tgcacccaag ttaggagaaa aacatatggt aagtgttaat cgtgggagcag
3051 tgtggcgaat cgcgaattgg gtttggcact tagatttcga tggcgctaga
3101 gacgccgttg gcgcgagcac catcgacctc atttttatgc gcgtgggaca
3151 ttgctgcaag agttttgagc atcgaatccc gcgtcgac
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